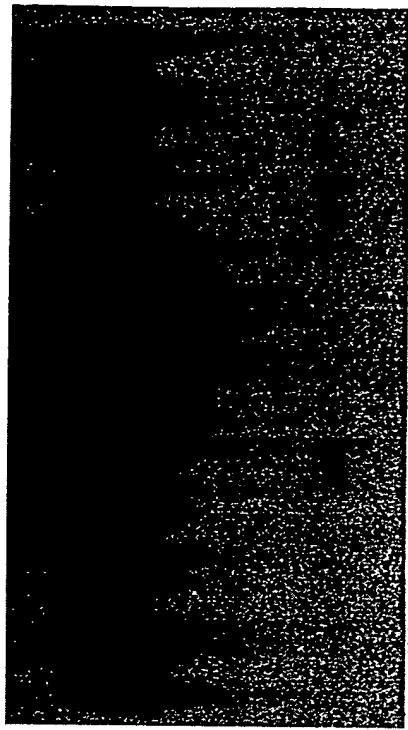


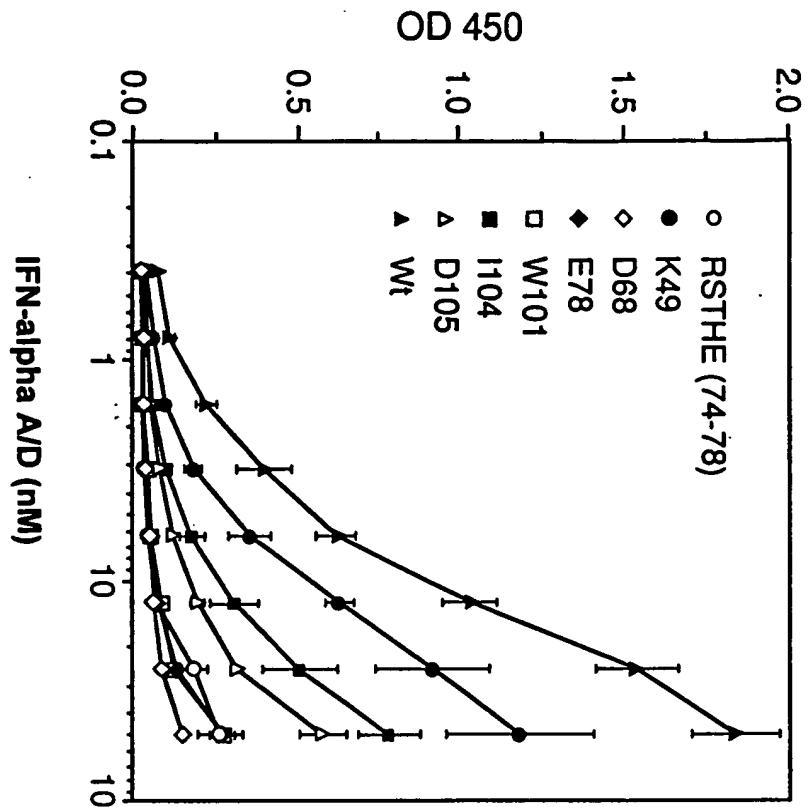
**Figure 1**

$\alpha$ IFN-2 mAb ( $\mu$ g/ml)	-	+	+	+	+	+	+	+	+	+	+
IgG	-	-	9F3	3B7	3B7	1D3	1D3	1F3	1F3	-	-
1	1	1	10	1	10	1	10	1	10	-	-

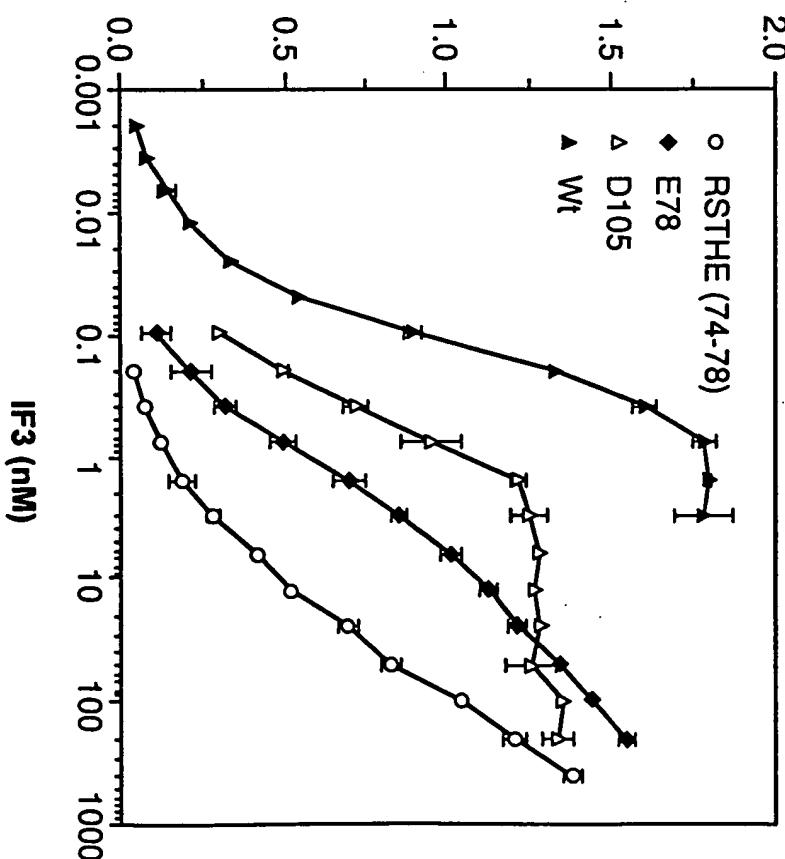
IGSF -



**Figure 2**

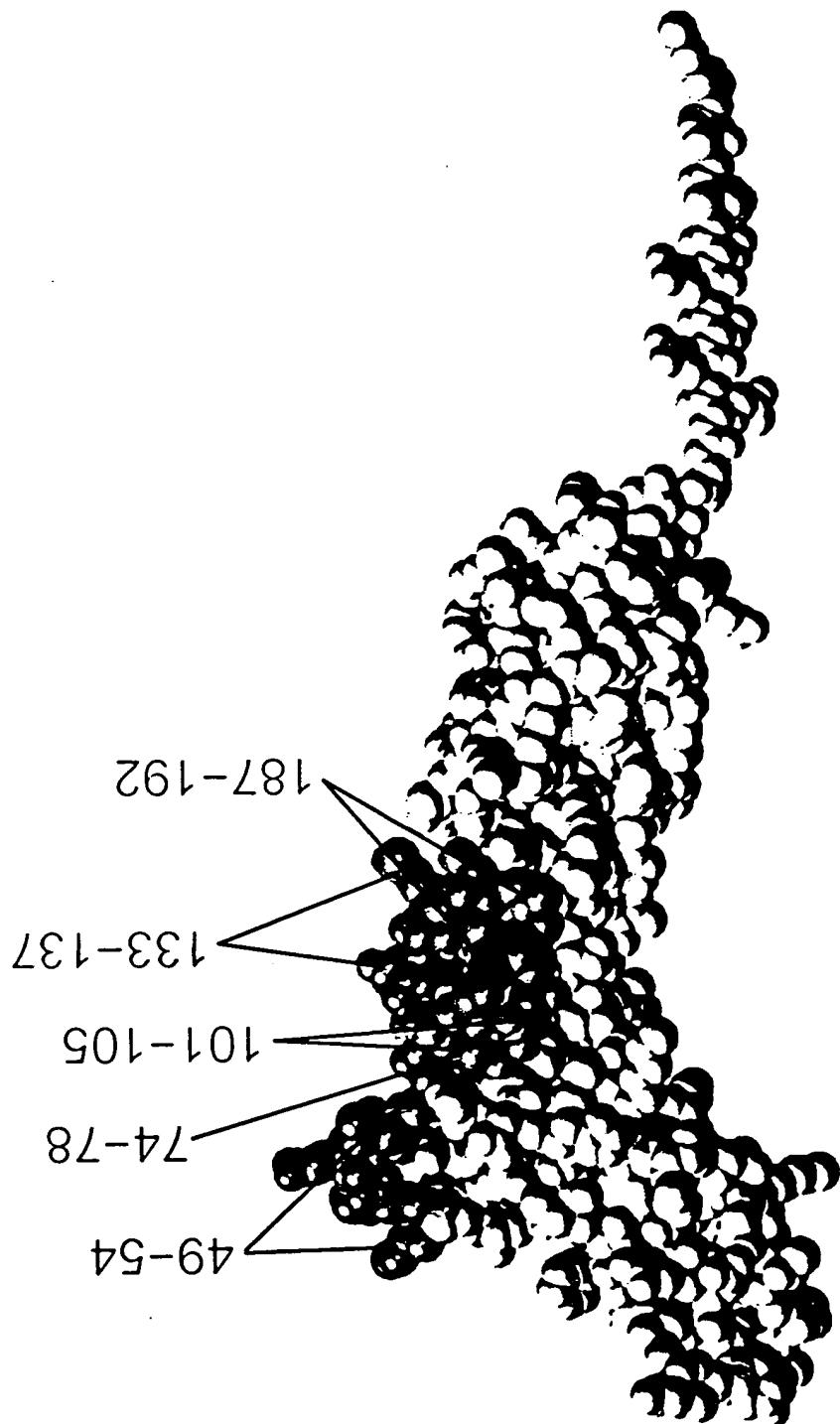


**Figure 3A**



**Figure 3B**

Figure 4



1 GAATTCTAA AATAGCMA GATGCTTG AGCCAGATG CCTCATCGT CAGACTT AATGGTT TCATGGTGA TATCAGCCC GTGTTGGTA  
CTAAGGATT TTATCGTT CTACGAAAC TCGGTCTAC GGAGTAGCA GTCTAGTGA TAAACCAAG AGTACCACT ATAGTCGGAG CACAAACCAT  
Ile

human alpha beta receptor

101 TTTCATATGA TTCGCCGTGAT TACACAGATG AATCTGCAC TTCAAGATA TCATGCAGA ATTCCGGTC CATTTATCA TGGGAATAA AAAACACTC  
AAGTATACT AAGCGGACTA ATGTGTCAC TTAGAACGTG AAAGTCTAT AGTACGCTT TAAAGGCCAG GTAGAATAGT ACCCTTATT TTTGGTGAG  
2 SerTyras pSerProAsp TyrThrAsp Tyro3Lck AspPleuLeu SerLeuWarg SerLeuWarg snPheArgse rileuseR TrpGluleuLysAsnHisSer

201 CATGTACCA ACTCACTATA CATGCTGTA TACATCATG AGTAAACCAAG AAGATTGAA GTGGTTAAG AACTGTGCAAG ATACCAAG ATCATTTGT  
GTAACATGGT TGAGCTATAT GTAACGACAT ATGTTAGTAC TCATTGGTC TTCTAACTT CCACCAATC TTGACACGTT TATGGTGTGTC TAGTAAACCA  
35 IleValPro ThrHistYrt hrLeuLeuTy rThrIleMet SerLysPro LuAspLeuLys sValVallys AsnCysAla snThrThrR gserPheCys

301 GACCTCACAG ATGAGTGGAG AACACACAC GAGGCCATAG TCACCGTCT AGAGGATC AGCGGACA CAACGTGTT CAGTGTCA CACAATTCT  
CTGGAGTGTCT TACTCACCTC TTGGTGTG CTCCGGATAC AGTGGCAGGA TCTTCCTAAG TCGCCTGT GTGCAACAA GTCAACGAGT GTGTTAAAGA  
68 AspPleuThra spGluTrpAsp gserThrHis GluAlaTyv alThrValle uGluGlyphe SerGlyAsnR hrThrLeuPhe eSerCysSer HisAsnPhetRp

401 GGCTGGCCAT AGACATGCT TTGAAACCAC CAGAGTTGA GATTGGTGT TTACCAACC ACATTATGT GATGGTGAAT TTCCATCTA TTGTTGAGGA  
CCGACCGGTA TCTGTACAGA AAACCTGGTG GTCTCAAATC CTAACAAACCA AAATGGTGG TGAATTACA CTACCACTT AAAGGTAGT AACAACTCCT  
102 LeuAlaIle eAspMetSer PheGluProP roGluPheGl uIleValGly Phethrash isileAsnVa 1MetVallys PheProSerI leValGluglu

501 AGAATTACAG TTGATTAT CTCTCGTCAT TGAAGAACAG TCAGAGGGAA TTGTTAAGAA GCATAAACCA GAAATAAAG GAAACATGAG TGGAAATTCTC  
TCTTAATGTC AAACTAATA GAGAGCAGTA ACTCTGTC AGTCTCCCT AACATTCT CGTATTGGG CTTATTTCTT CTTGTACTC ACCTTAAAG  
135 GluLeuGln PheAspLeu erLeuValIle eGluGluGln SerGluGlyI leValLysLys ShisLysPro GluIleLysG lyAsnMetSe rGlyAsnPh

601 ACCTATATCA TTGACAAAGTT ATTCCAAAC AGCAACTACT GTGTATCTGT TTATTAGAG CACAGTGTG AGCAAGCAGT AATAAGTCT CCCTAAAT  
TGGATATAGT AACTGTCAA TTAGGTTG TGCTGTGTA CACATAGACA AATAAATCTC GTGTCACTAC TCGTCTGTC TTATTCTAGA GGAATTAA  
701 GCAACCTCTT TCCACCTGGC CAGGAATCAG AATCAGCAGA ATCTGCCAC AAACTCACA CATGCCACC GTGCCAGCA CCTGAACCTC TGGGGGACCC  
CGTGGGAGGA AGGTGGACCG GTCTCTAGTC TTAGTCGTCT TAGACGGCTG TTGAGGTGT GTACGGTGG CACGGTGT GGACTGAGG ACCCCCTGG  
202 ThrLeuLeu upProProGly GluGluSerG luserAlaAsp LysThrHist hrCysProP roCysProAla ProGluLeu euglyGlyPro  
19G1

801 GTCAGTCTTC CTCTCCCC CAAACCCAA GGACACCTC ATGATCTCC GGACCCCTGA GGTACACATGC GTGGGGTGG ACGTGAGCCA CGAAGACCT  
CAGTCAGAAG GAGAAGGGG GTTTGGGTT CCTGGGAGG TACTAGAGG CCTGGGACT CCAGTGTACG CACCACCC TGCACTCGGT GCTCTGGGA  
235 SerValPhe LeuPheProP roLysProLys sAspThrLeu MetIleSerA rgThrProG1 uValThrCys ValValVala spValSerH1 sGluAspro

Figure 5A

## Figure 5B

901 GAGGTCAGT TCAACTGGTA CGTGGACCGC GTGGAGGTGC ATAATGCCAA GACAAGCCG CGGGAGGAGC AGTACAACAG CACGTCACGA GTGGTCAGGC  
CTCCAGTTCA AGTGACCAT GCACCTGCCG CACCTCCACG TATTACGGTT CTGTRTCGGC GCCCTCCTCG TCATGTGTC GTGCATGGCT CACCAGTCGC  
268 GluValLysPheAsnTrpTy ValAspGly ValGluValH isAsnAlaLysThrLysPro ArgGlugIuLysInTrpAsnSe rThrTyRarg ValValSerVal  
1001 TCCTCACCGT CCTGCCACAG GACTGGCTGA ATGGCAAGGA GTACAGTC AAGGTCTCCA ACAAGCCCT CCCAGCCCC ATCGAGAAA CCATCTCCAA  
AGGAGTGGCA GGACGTRGGC CTGACCGACT TACGGTTCT CATGTCACG TTCCAGAGGT TGTTGGGA GGGTCGGGG TAGCTCTTT GGTAGAGGTT  
302 LeuThrVal lleuHisGln AspTrpLeu snglyLysAl uTyryLysCys LysValSerA snLysAlaLys uproAlaPro ileGluLysT hrileSerLys  
335 AlalysGly GlnProArgG luproGlnVa lTyryThrLeu ProProSerA rgGluGluMe tThrLysAsn GlnValSerL eurhrcysLe wValLysGly  
1201 TTCTATCCA GCGACATCGC CGTGGAGTGG GAGAGCAATG GGCAGCCGA GAACAACATC AAGACCAACGC CTCCTGTCTT GGACTCCGAC GGCTCCTCT  
AAGATAGGGT CGCTGTAGCG GCACCTCAC CTCTCGTTAC CCGTCGGCT CTGTGTGATG TTCTGGTGCG CCGAGGCTG CCGAGGAAGA  
368 PheTyrProSer erAspIleAl aValGluTrp GluSerAsnG lyGlnProG l uAsnAsnTyr LysThrThrP reproValle uAspSerAsp GlySerPhePhe  
1301 TCCTCTACAG CAAGCTCACC GTGGACAAGA GCAGGTGGCA GCACGGGAAC GTCTTCTCAT GCTCCGTAT GCTCCGTAT GCTATGGCT CTGACACAGCA  
AGGAGATGTC GTCAGTGG CACCTGTCT CGTCACCGT CGTCCCCTTG CAGGAGAGTA CGAGGCACTA CGTACTCCGA GACGTGTTG TGATGTGCGT  
402 LeuTyrSe rLysLeuThr ValAspLysS erArgTrpGlnGlnGlyAsn ValPheserC ysSerValMe thIsGluVala LeuHisAsn iStyryThrGln  
1401 GAAGAGCCTC TCCCTGTCTC CGGGTAATG AGTGGACGG CCCTAGAGTC GACCTGCAGA AGCTTAGAAC CGAGGGCCG CCATGGCCCA ACTGTGTTAT  
CTTCTCGGAG AGGGACAGAG GCCCATTAC TCACGCTGCC GGGATCTCAG CTGGACGTCT TCGAATCTG GCTCCCCGGC GGTACCCGGT TGAACAAATA  
435 LyserSerLeu SerLeuSerP roGlyLysP \* (SEQ ID NO. 26) sv40 early  
  
poly A

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ACGTGAAATA TTACCAATGT TTATTTCGTT ATCGTAGTGT TAAAGTGT TATTTCGAA AAAAGGAC GTAAAGTCAA CACCAACAG GTTGTGAGTAG  
1601 AATGTATCTT ATCATGCTG GATCGATCGG GAATAATC GCGCGACAC CATGGCTGA AATAACCTCT GAAGAGGAA CTGGTTAGG TACCTCTGA  
TTACATAGAA TAGTACAGAC CTAGCTAGCC CTTAATAG CGCGCTGTG GTACCGGACT TTATGGAGA CTTCTCCT GAACCAATCC ATGGAAGACT  
sv40 origin

1701 GGGGAAAGA ACCAGCTGTG GAAATGTGTG CAGTAGGGT GTGGAAAGTC CCCAGGCTCC CGAGCAGGCA GAAGTATGCA AAGCATGCA CTCATATTG  
CCGCCTTCTT TGGTGCACAC CTTACACACA GTCAATCCCA CACCTTCAG GGGTCCGAGG GGTGTCCTG CTCATACGGT TTCTGACGTA GAGTTAATCA  
1801 CAGCAACCAAG GTGTGAAG TCCCCAGGCT CCCAGCAGG CAGAAGTATG CAAGCATGC ATCTCAATTA GTCAAGCAACC ATAGTCCCCG CCCTAACTCC  
GTCGTTGGTC CACACCTTC AGGGTCCCA GGGGTGTC GTCTTCATAC GTTCTGTAAG CAGTCGTGG TATCAGGGCG GGGATTGAGG

1901 GCCCATCCCC CCCCTAATC CGCCCAAGTC CGCCCATCT CGCCCCATG GCTACTAAT TTTTTATT TATGCAGAGG CCGAGCCGC CTCGGCCTCT  
CGGCTAGGGC GGGATGAG GGGGTCAAG GGGGTAAGA GGCGGGTAC CGACTGATA AAAAATAA ATACGTCTCC GGCTCCGGCG GAGCGGAGA  
2001 GAGCTATTC AGAAGTAGTG AGGAGGTT TTTGAGGCC TAGGTTTG CAAAAGCTG TAAACAGCTT GCAACTGGCC GTCGTTTAC AACGTCGTA  
CTCGATAAGG TCTTCATCAC TCTCCGAAA AACCTCCGG ATCCGAAAC GTTTCGAC AATTGTCGAA CGGTGACCGG CAGCAAATG TTGCGAGCT  
start pUC118

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GACCCCTTG GGACCCAAT GGGTGAATT AGCGAACGT CGTAGGGG GGAAGGGTC GACCGCATTA TCGCTTCTCC GGGCGTGGCT AGCGGAAAGG  
2201 CACAGTTGC GTAGCCTGAA TGGCGAATGG CGCCGTGATGC GGTATTCT CCTACGCAT CTGTCGGTA TTTCACACCG CATACTCAA AGCAACCATA  
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2301 GTACGCGCCC TGTAGGGCG CATTAGCGC GGCGGGTGTG GTGGTTACGC GCAGCGTGC CGCTACACTT GCCAGGCC TAGCGCCCGC TCCTTCGCT  
CATGGGGG ACATGGCGC GTAATTGCG CGGCCACAC CACCAATGCG CGTCGCACTG GCGATGTGAA CGTGCACGGG ATCGGGCG AGGAAGCGA  
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ACCTGAGAAC AAGGTTGAC CTTGTTGTA GTGGGATAG AGCCCGATAA GAAACTAA TATCCCTAA AACGGCTAA GCGGATAAC CAATTGTTA  
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3001 AGGGCTCGT GATACGCTA TTTTATAGG TAAATGTCAT GATAATAATG GTTCTTCTAGA CGTCAGGTGG CACTTTCGG GGAATGTC GGGGAACCC  
TCCCGAGCA CTATCGGAT AAAATATCC ATTACAGTA CTATTATAC CAAAGAATCT GCACTCACC GTGAAAGCC CCTTACACG CGCCTGGGG

Figure 5C

3101 TATTGTGTTA TTTTCTAA TACATCAA TATGTATCCG CTCATGAGAC AATAACCTG ATAATGCTT CAATAATT GAAAAGGA GAGTATGAGT  
 ATAACAAT AAAAGATT ATGTAAGTT ATACATAGGC GAGTACTCTG TTATGGAC TATTACGAA GTTATTATAA CTTTTCCCTT CTCATACTCA  
  
 3201 ATTCAACATT TCCGTGTCGC CCTTATCCC TTTTGCGG CATTTGCCT TCCTGTTT GTCACCCAG AACGCTGGT GAAAGTAAA GATGCTGAAG  
 TAAGTGTAA AGGCACAGGG GGAATAAGG AAAAACGCC GTAAAGGA AGGCAAAAA CGAGTGGTC TTGCGACCA CTTCTCATTT CTACGACTTC  
  
 3301 ATCAGTGGG TGCACGGGT GCTTACATCG AACTGGATCT CAACAGGGT AAGATCCTG AGAGTTTCG CCCCCGAAGAA CGTTTCCAA TGATGAGCAC  
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 3601 TTCTGACAC GATGGAGGA CGAAGGGAGC TAACCAGCTT TTGACACAC ATGGGGATC ATGTAACTCG CCTTGATCGT TGGAAACCGG AGCTGAATGA  
 AAGACTGTG CTAGCCTCT GGCTTCCTCG ATTGCGAA AACCGTGTG TACCCCTAG TACATGAGC GGAACTAGCA ACCCTGGCC TCGACTTACT  
  
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 3901 GTGAGCGCTG GTCTCGCGGT ATCATGCG CACTGGGCC AGATGTAAG CCCTCCGTA TCGTAGTTAT CTACAGCAG GGGACTCAGG CAACTATGAA  
 CACTCGCACC CAGAGCGCCA TAGTAACGTC GTGACCCCCGG CCTACATTG GGGAGGGCAT AGCATCAA GATGTCG CCGCTAGTCC GTTGATACCT  
  
 4011 TGAAACGAAT AGACAGATCG CTGAGATAGG TGCCCTACTG GACTCTATCC ACGGAGTGC TAATCGTAA CCATGACAG TCTGGTCAA ATGAGTATAT ATGAATCTA ACTAAATT  
 ACTGTGTTA TCTGCTTAGC GACTCTATCC ACGGAGTGC TAATCGTAA CCATGACAG TCTGGTCAA ATGAGTATAT ATGAATCTA ACTAAATT  
  
 4101 CTTCATTT AATTAAAG GATCTAGGTG AAGATCCTT TTGATAATCT CATGACCAA ATCCCTAAC GTGAGTTTC GTCCACTGA GCGTCAGACC  
 GAAGTAAAAA TTAAATTTC CTAGATCCAC TTCTAGGAA AACTATTAGA GTACTGGTT TAGGAAATTG CACTCAAAG CAAGGTGACT CGCAGTCTG  
  
 4201 CGGTAGAAA GATCAAAGA TCTCTTGAG ATCCCTTT TCTCGCGTA ATCTGCTGCT TGCACACAA AAAACACCG CTACAGCGG TGGTTGTT  
 GGATCTTT CTAGTTCCCT AGAAGAACTC TAGGAAAGA AGACGCGCAT TAGACGACGA ACGTTGTTT TTGCGTGC GATGCGGCC ACCAACAAA  
  
 4301 GCCGGATCAA GAGCTACCA CTCCTTCC GAAGGTAAC GGCTTCAGCA GAGGCGAGAT ACCAATACT GTCTCTAG TGTAGCCGTA GTTGGCCAC  
 CGGCCTAGTT CTCGATGGTT GAGAAAAGG CTTCATGA CGAAGTCGT CTCGCGTCA TGGTTATGA CAGGAAGATC ACATGGCAT CAATCCGGTG

**Figure 5D**

4401 CACTTCAGA ACTCTGTAGC ACCGCCTACA TACCTCGCTC TGCTAATCCT GTCACCGAGTG GCTGCTCCA GTGGCGATAA GTCGTGCTT ACCGGGTTGG  
 GTGAAGTCT TGAGACATCG TGGGGATGT ATGGAGCGAG ACGATTAGGA CAATGGTCAC CGACGACGGT CACCGCTATT CAGCACAGAA TGGCCCAAC  
 4501 ACTCAAGAGC ATAGTACCG GATAAGGGC AGCGCTGGG CTGAACGGG GGTTCGTGCA CACAGCCCAG CTGGAGCGA ACGACCTACA CGAACACTGAG  
 TGAGTCTGCC TATCAATGGC CTATCCGGG TCGCAGCCC GACTGCCCC CCAAGCACGT GTGCGGTC GAACTCGCT TGCTGGATGT GGCTGAATC  
 4601 ATACCTACAG CGTGAGCATT GAGAAAGGC CACGCTCCC GAAGGGAGAA AGGGGACAG GTATCGGTA AGCGGCAGG TCGGACAGG AGAGCGCAC  
 TATGGATGTC GCACTCGTAA CTCTTCGGG GTGCGAAGGG CTCCTCTT TCGCGCTGTC CATAGGCCAT TCGCCGTCCC AGCCTGTC TCTCGCGTGC  
 4701 AGGGAGCTTC CAGGGGAAA CGCCTGGTAT CTTATAGTC CTGTCGGTT TCGCCACCTC TGACTTGAGC GTCGATTGTT GTGATGCTCG TCAGGGGGC  
 TCCCTCGAAG GTCCCCCTT GCGGACATA GAAATATCAG GACAGCCAA AGCGGTGGAG ACTGAACTCG CAGCTAAAG CACTACGAGC AGTCCCCCG  
 4801 GGAGCCTATG GAAAACGCC AGCAACCGGG CCTTTTACG GTTCTG GCC TTTGCTGCC CTTTGCTCA CATGTTCTT CCTGGTTAT CCCGTATC  
 CCTCGGATAC CTTTTGCGG TCGTTCGCC GAAAAATGC CAAGGCCGG AAACGACCG GAAAAGGT GTACAAGAAA GGACGCCATA GGGACTAAG  
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 ACACCTATTG GCATAATGGC GAAACTCAC TCGACTATGG CGAGGGCGT CGGCTGCTG GCTCGCTCG CTCAGTCAGT CGCTCTCG CTTCTCGCG  
 5001 CCAATACGCA AACCGCCCT CCCCCGGCGT TGGCGATT ATTAAATCAG CTGGCACGAC AGGTTCCCG ACTGGAAGC GGGCAGTGAG CGCAACGCAA  
 GGTATGCGT TTGGGGAGA GGGGGCGCA ACCGGCTAG TAATTAGTC GACCGTGCTG TCCAAGGGC TGACCTTGC CCCGTCACTC CGGTGCGTT  
 5101 TTAATGTGAG TTACCTCACT CATTAGGCAC CCCAGGCTT ACACTTATG CTTCCGCTC GTATGTGTC TGAATGTG AGGGATAAC AATTCAAC  
 AATTACACTC AATGGGTGA GTAATCCGGT GGGTCCGAAA TGTGAAATAC GAAGGGGAG CATAACAC ACCTTAACAC TCGCTTATG TAAATGTG  
 5201 AGGAAACAGC TATGACCATG ATTACGAATT AATTCGAGCT CGCCCGACAT TGATTATGA CTAGTTATA ATAGTAATCA ATTACGGGT CATTAGTC  
 TCTTTGTGCG ATACTGGTAC TAATGCTAA TTAAGCTGA GCGGGCTGTA ACTAAATACT GATCAATAAT TATCATAGT TAATCCCCA GTAAATCAAGT  
 from pPMLCMV beginning to HindII, enhancers and promoter

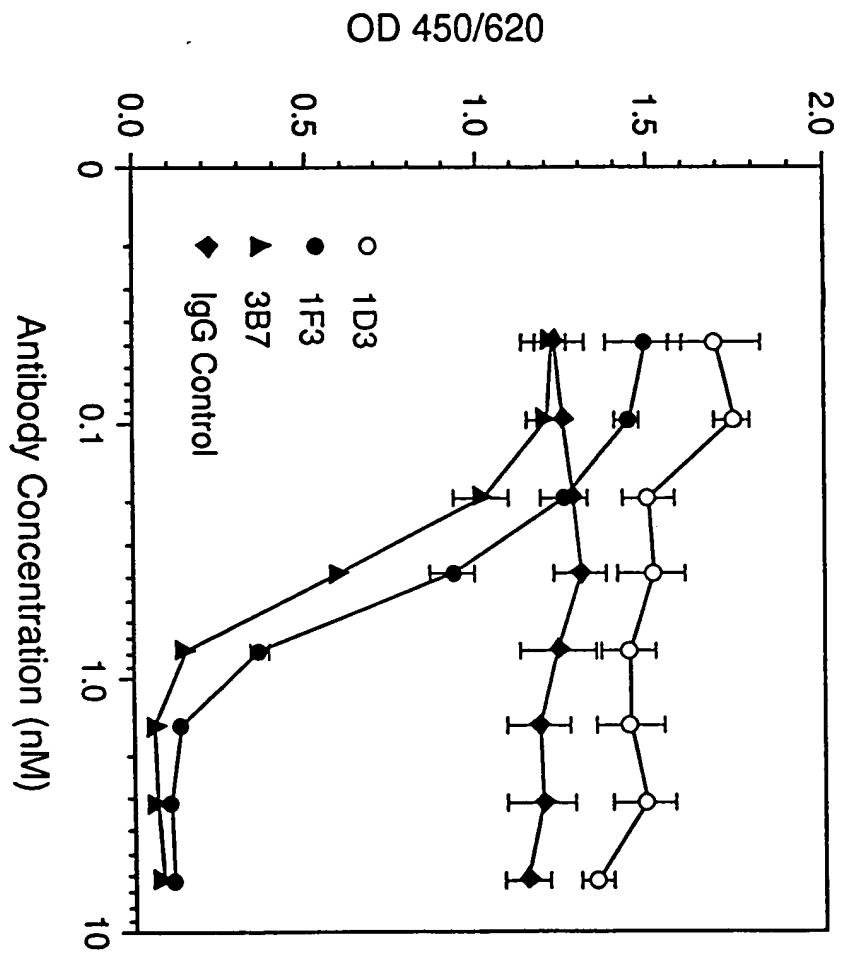
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 5501 CAAGTACGCC CCCTATTGAC GTCAATGAGG GTAATGGCC CGCCTGGCAT TATGCCAGT ACATGACCTT ATGGGACTT CCTACTTGCG AGTACATCTA  
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**Figure 5E**

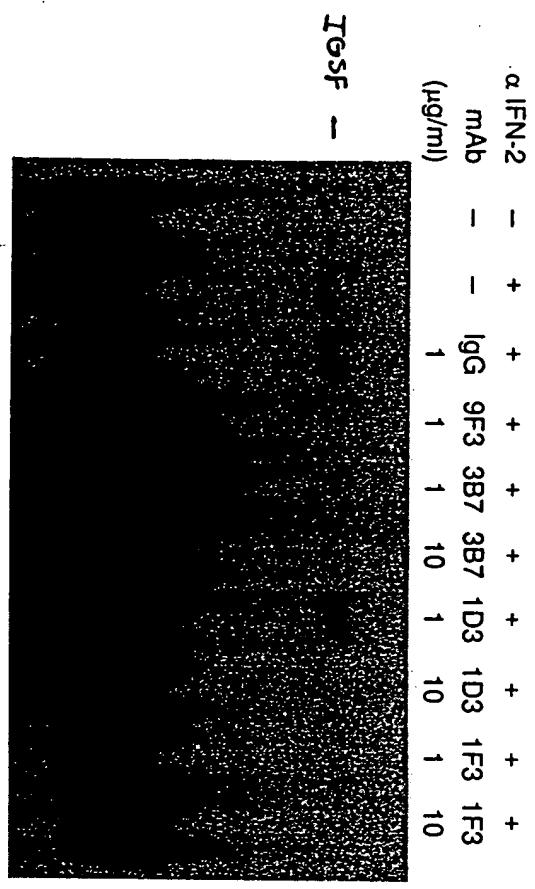
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5801 TACGGTGGGA GGTCTATATA ACCAGAGCTC GTTGTAGTGA CCGTCAGATC GCCTGGAGAC GCCATCCACG CTGTTTGAC CTCCATAGAA GACACCGGA  
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sp6 promoter  
sp6 RNA start

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cloning linker

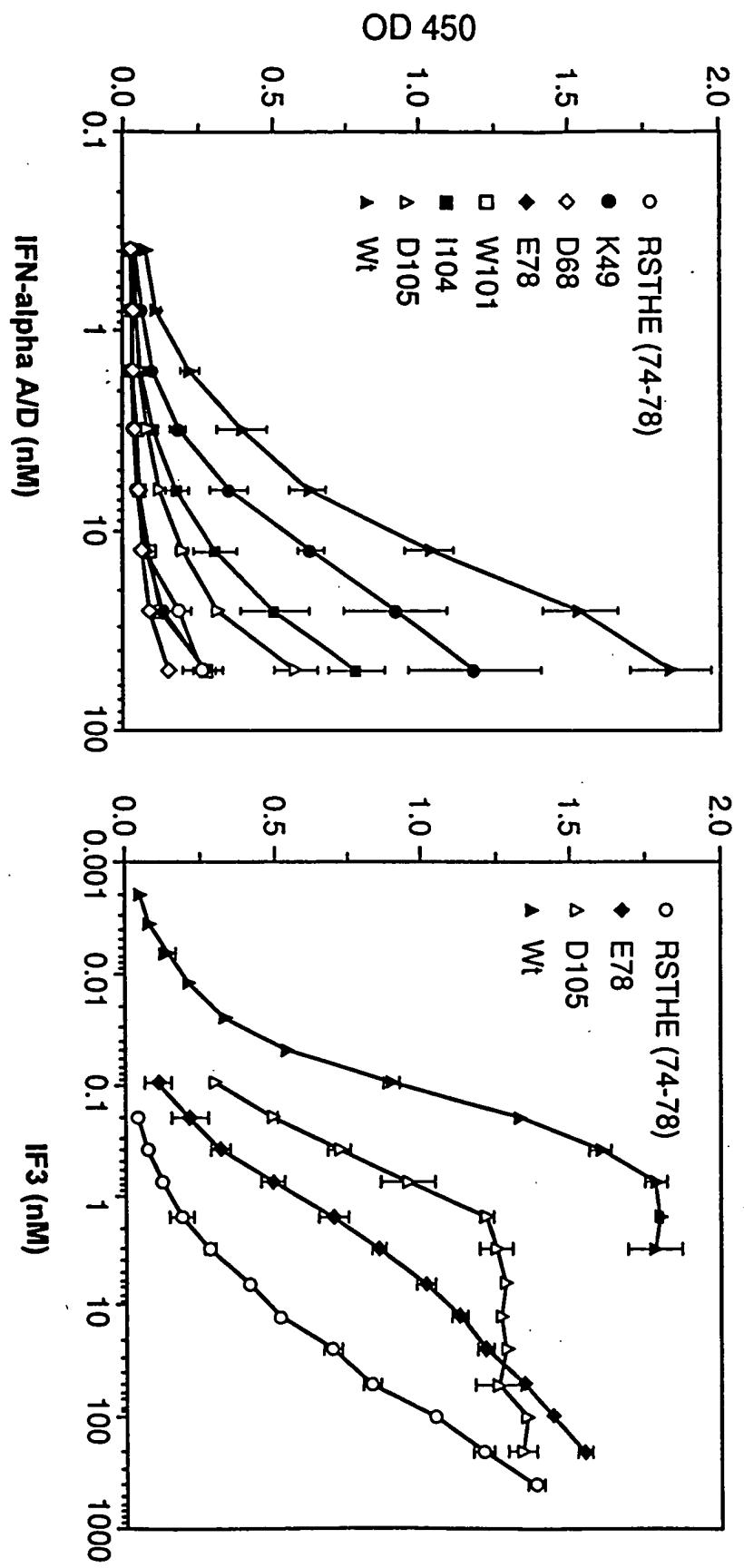
**Figure 5F**



**Figure 1**

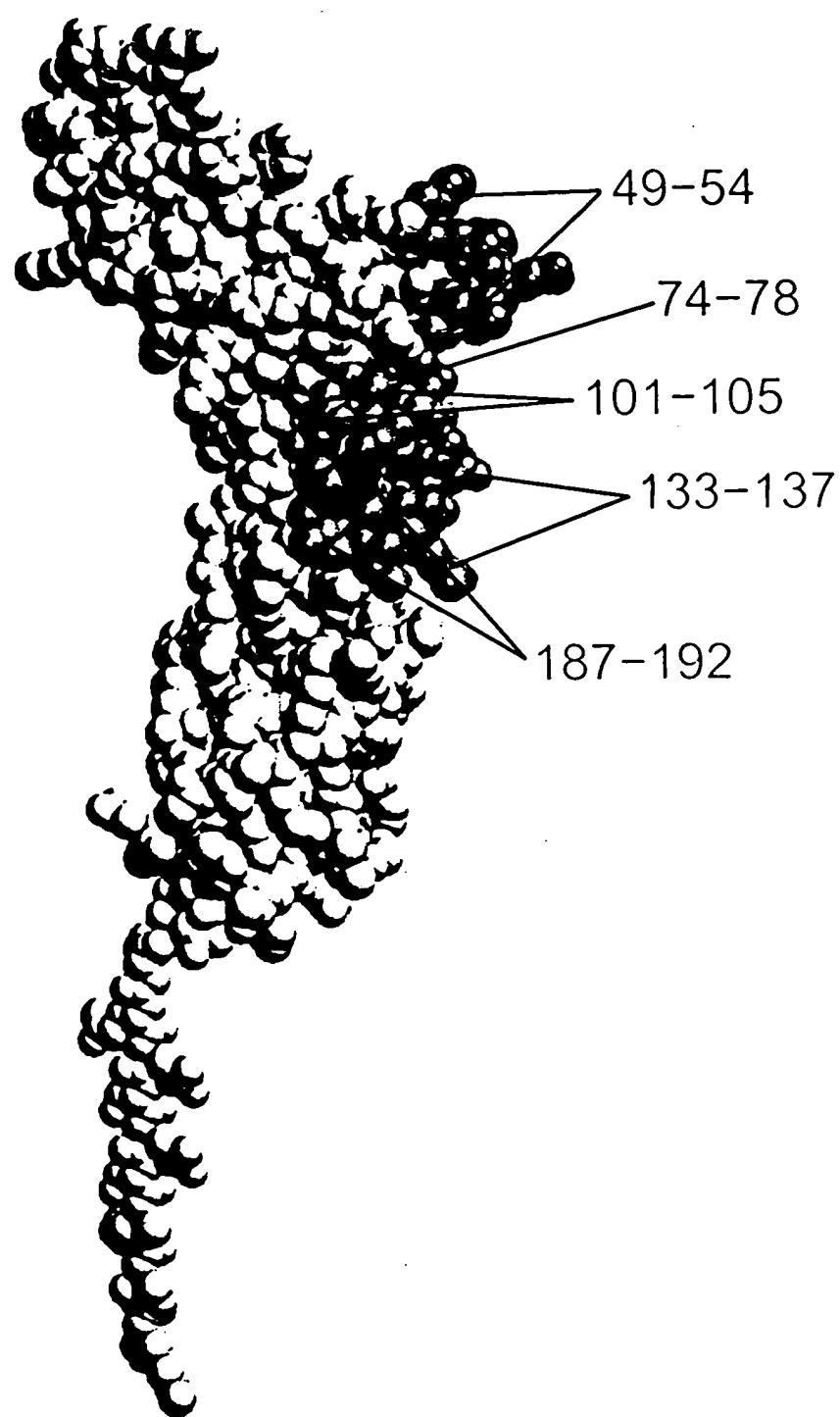


**Figure 2**



**Figure 3A**

**Figure 3B**



**Figure 4**

1 GAATTCTAA AATAGCAA GATGCTTG AGCCAGATG CCTTCATCGT CAGATCACTT AATTGGTC TCATGGTGA TATCAGGCTC GTGTGGTA  
CTAAGGATT TTATCGTT CTACGAAAC TCGTCTAC GGAGTAGCA GTCTAGGAA TAAACCAAG AGTACCACT ATAGTCGGAG CACAACCAT  
Ile

human alpha beta receptor

101 TTTCATATGA TTCGCCGTAT TACACAGATG AATCTGCAC TTCAAGATA TCATTCGAA ATTCCGGTC CATCTTATCA TGGAAATAA AAAACCACTC  
AAGTATACT AAGCGGACTA ATCTGTCTAC TTAGAACGTG AAAGTTCTAT AGTACGCTT TAAAGGCCAG GTAGAATAGT ACCCTTATT TTTGGTGAG  
2 SERTYras pSerProasp TyrThrAsp IuserCysth rPhelysile SerLeuWargA snPheArgSe rIleLeuSer rPgluLeuLysAshKisser

201 CATGTACCA ACTCACTATA CATTGCTGTA TACATCATG AGTAAACCAAG AAGATTGAA GGTGGTTAG AACTGTGAA ATACCAAG ATCATTGTT  
GTACATGGT TGAGTGTAT GTAACGACAT ATGTTAGTAC TCATTGGTC TTCTAACTT CCACCAARTC TTGACACGTT TATGGTGTTC TAGTAACAA  
35 IleValPro ThrHistYrt hrLeuLeuTy rThrIleMet SerLysProg Luaspleuly sValVallys AsnCysAlaA snThrThrR gSerPheCys

301 GACCTCACAG ATGAGTGGAG AACACACAC GAGGCTATG TCACCGTCT AGAAGGATC AGCGGGAA CAACGTTGTT CAGTGTCA CACAATTCT  
CTGGAGTGTCT TACTCACCTC TTCTGTGTG CTCCGGATAC AGTGGCAGGA TCTTCCTAAG TCGCCTGT GTGCAACAA GTCAACAGGT GTGTTAAAGA  
68 Aspleuthra spGLUTpar gserThrHis GluAlaTyrv alThrValle uGluLysPhe SerGlyAsn hrThrLeuPh eserCysSer HisAsnPhetTP

401 GGCTGGCCAT AGACATGCT TTGAAACCAC CAGAGTTGA GATGTTGTT TTACCAACC ACATTAATGT GATGGTGAAT TTCCATCA TTGTTGAGGA  
CCGACCGGTA TCTGTACAGA AAATCTGGTG GTCTCAAAT CTAAACCA AAATGGTTG TGAATTAACA CTACCACTT AAAGGTAGT ACAACTCCT  
102 LeuAlaIle eAspMetSer PheGluProg roGluPheGL uIleValGly PheThrAsnH isileAsnVa 1MetVallys PheProSerI levalGluGlu

135 501 AGAATTACAG TTGATTAT CTCTCGTCAT TGAGAACAG TCAGAGGAA TTGTTAGAA GCATAACCC GAAATAAAG GAAACATGAG TTGAAATTG  
TCTTAATGTC AAACTAATA GAGAGCAGTA ACTCTGTCT AGTCTCCCCT AACATTCT CGTATTGGG CTTTATTTC CTTGTACTC ACCTTAAG  
GlueLeugln PheAspleus erleuValIle eGluGluGln SerGluGlyI leValLysly shisLysPro GluIleLysG lyAsnMetSe rGlyAsnPh

601 501 ACCTATATCA TGACAAAGT ATTCCAAAC AGCAACTACT GTGTATCTGT TTATTTAGAG CACAGTGT AGCAAGCAGT AATAAGTCT CCCTAAAT  
TGGATATAGT AACTGTCAA TTAGGTGT TGCTGTGTA CACATAGACA ATAATCTC GTGTCACTAC TCGTGTGCA TTATTCAGA GGGAAATTAA  
j8 ThrTyrIleI leAspLysle uIleProAsn ThrAsnTyrc ysValServa lTyrLeuGlu HisSerAspI lugInAlaVa lIleLySseR ProLeuLysCys

701 501 GCACCCCTCT TCCACCTGGC CAGGAATCAG AATCAGCAG ATCTGCCAC AAAACTCACA CATGCCACC GTGCCAGCA CCTGAACCTC TGGGGGACCA  
CGTGGGAGGA AGGTGGACCG GTCTCTAGTC TTAGTCGTCT TAGACGGCTG TTGTGAGTGT GTACGGTGG CACGGTGT GGACTTGAGG ACCCCCTGG  
202 ThrLeuIle uproProGly GluGluSerG IuserAlaIle userAlaAsp LysThrHist hrCysProPr oCysProAla ProGluLeuL euGlyGlyPro  
19G1

801 501 GTCAAGTCTTC CTCTCCCCC CAAACCAA GGACACCCCTC ATGATCTCC GGACCCCTGA GGTCACTGC GTGGTGGTGG ACGTGAGCCA CGAAGACCC  
CAGTCAGAAG GAGAAGGGGG GRTTGGGTT CCTGTGGGAG TACTAGAGG CCTGGGACT CCAGTGTACG CACCAACCC TGCACCTCGGT GCTTCCTGG  
235 servAlphe LeupheProP rolysProL yAspThrLeu MetIleSerA rgThrProG1 uValThrCys ValValVala spValSerhi sGluasPro

Figure 5A

901 GAGGTCAGT TCAACTGGTA CGTGGACGCC GTGGAGGTGC ATAATGCCAA GACAAGCCG CGGGAGGAGC AGTACAACAG CACGTACCGA GTGGTCAGGC  
 CTCCAGTTCA AGTGACCAT GCACCTGCCG CACCTCCAG TATTACGGTT CTGTTTCGGC GCCTCTCTCG TCATGTTGTC GTGCATGGCT CACCAAGTCGC  
 268 GluValLysP heAsnTrpTy rValAspGly ValGluValH isAsnAlaLy sThrLysPro ArgGluglu InTrYrAsnE rThrTyrArg ValValServal  
 1001 TCTTCACCGT CCTGCACCAAG GACTGGCTGA ATGGCAAGGA GTACAAGTC ACAGGCCCCT CCCAGCCCCC ATCGAGAAGA CACATCTCAA  
 AGGAGTGGCA GGACGTTGGTC CTGACCCGACT TACCGTTCT CATGTTACG TTCCAGAGGT TGTTGGGA GGGTGGGG TAGCTCTTT GTAGAGGTT  
 302 Leuthrva lleuHisGln AspTrpLeu snglyLysAl uTyrLysCys LysValSerA snlysAlaLe uproAlaPro ilegulyst hrleSerLys  
 335 AlalysGly GlnProArgg IuproGlnVa lTyrThrLeu ProProSerA rggIuGlume tThrLysAsn GlnValSerL euthrCysLe uValLysGly  
 301 TTCTATCCCA GCGACATCGC CGTGGAGTGG GAGAGCAATG GGCAGCCGA GAACAACTAC AAGACCACGC CTCCCGTGT GGACTCCGAC GGCTCCTCT  
 AGGATAGGGT CGCTGTAGCG GCACCTCACCT CTCTCGTAC CGTCCGGGCT CTGTTGATG TTCTGGTGC GAGGGCACGA CCTGAGGCTG CCGAGGAAGA  
 368 PheTyrProS erAspIleAl avAlglutrp GluSerAsnG LysThrThrP reproValle uAspSerAsp GlySerPhePhe  
 1301 TCCTCTACAG CAAGCTCACC GTGGACAAGA GCAGGGCA GCAGGGGAAC GTCTTCTCAT GCTCCGGTAT GCTCCGGTAT GCTGAGGCT CTGCACAAACC ACTACACGCA  
 AGGAGATGTC GTCGAGTGG CACCTGTTCT CGTCCACCGT CGTCCCGCT CGTCCCGCT CAGAAGAGTA CGAGGCACTA CGTACTCCGA GACGTTGTTG TGATGTCGCT  
 402 LeutYrSe rLysLeuthr ValAspLysS erArgTrpGln gLnglyAsn ValPheSerC ysSerValMe thiGluVala LeuHiASnH iStYrThrGln  
 1401 GAAGAGCCTC TCCCTGTC CCGGTAATG AGTGGACGG CCCTAGAGTC GACCTGCAGA AGCTTAGAAC CGAGGGCCG CCATGCCA ACTGTTAT  
 CTCTCGGGAG AGGGACAGAG GCCCATTAC TCACCTGCC GGGATCTCAG CTGGACGTCT TCGAATCTG GCTCCCGGC GGTACCGGGT TGAACAAATA  
 435 LysserLeu SerLeuSerP roGlyLysP \* (SEQ ID NO.26) SV40 early  
 poly A

1501 TGCAGCTTAT AATGGTTACA ATAAGCAA TAGCATCACA AATTCAAA ATAAGCATT TTTTCACTG CATTCTAGT GTGGTGTGC CAAACTCATC  
 ACGTCGAATA TTACCAATGT TTATTTCGTT ATCGTAGTGT TAAAGTGT TATTTCGAA AAAAGTGAC GTAAGATCAA CACCAACAG GTTGTGAGTAG  
 2601 AATGTATCT ATCATGTCG GATCGATCGG GAATAATC GGCGCAGCAC CATGGCTGA ATAACCTCT GAAAGAGGA CTGGTTAGG TACCTCTGA  
 TTACATAGAA TAGTACAGAC CTAGCTAGCC CTCATTAAG CGCGTGTG GTACGGACT TTATGGAGA CTTCTCCTT GAAACCAATCC ATGGAAGACT  
 SV40 origin

1701 GGGCGGAAGA ACCAGGTGTG GAATGTGTG CAGTGTAGGT GTGGAAAGTC CCCAGGCTCC CCAGCAGGCA GAAAGTATGCA AAGCATGCA CTCATTAAGT  
 CCGCCTTCTT TGGTCGACAC CTTACACACA GTCAATCCCA CACCTTCAG GGGTCCGAGG GGTGTCCGT CTCATACGTT TCGTACGTA GAGTAATCA  
 1801 CAGCAACCAG GTGTGGAAAG TCCCCAGGCT CCCAGCAGG CAGAAGTATG CAAGCATGC ATCTCAATT GTCAAGCAACC ATAGCCCGC CCCTAACTCC  
 GTCGTGGTC CACACCTTC AGGGTCCCA GGGGTGTC GTCTTCATAC GTTGTGAG TAGAGTAAT CAGTCGTGG TATCAAGGGC GGGATTGGG

**Figure 5B**

1901 GCCCATCCCG CCCTAATC CGCCCAAGTC CGCCCCATG GCTACTAAT TTTTTTATT TATGCAGGG CCGAGGCCG CTCGGCCTCT  
CGGGTAGGGC GGGGATGAG GCGGTCAAG GCGGGAAGA GCGGGGTAC CGACTGATTA AAAAATAA ATACGTCTCC GGCTCCGGCG GAGCCGGAGA  
2001 GAGGTATCC AGAAGTAGTG AGGAGGCTT TTGGAGGCC TAGGTTTG CAAGACTG TAACAGCTT GGCACGGCC GTCGTTAAC AACGTGAGA CCTGAC  
CTCGATAAGG TCTTCATCAC TCCCTCGAAA AACCTCCCG ATCCGAAC GTTTCGAC AATGTGAA CGTGACCGG CAGCAATG TTGCAACT  
start pUC18

2101 CTGGGAAAC CCTGGGCTTA CCCAACTTAA TCGCCTGCA GCACATCCC CCTTCGCCAG CTGGGTAAT AGCGAAGGG CCCGCACCGA TGCCTCTCC  
GACCCTTTG GGACCGCAAT GCGTGAAATT AGCGAACGT CGTGTAGGG GGAAGGGTC GACCGCATTA TCGCTCTCC GGGGTGGCT ACCGGAAAGG  
~201 CAACAGTTGC GTAGCCTGAA TGGCGAATGG CGCCTGATGC GGTATTCT CCTACGCAT CTGCGGTA TTCACACCG CATACTCAA AGCAACCATA  
GTGTCGAACG CATCGACTT ACCGCTTACCG GCGGACTACCG CCATAAAGA GGAATGCGTA GACACCCAT AAATGTGGC GTATGAGTT TCGTGGTAT  
2301 GTACGCGCCC TGTAGGGCG CATTAGGCC GGGGGGTGTG GTGGTTACGC GCAGCGTAC CGCTACATT GCCAGGCC TAGGCCGC TCCTTCGCT  
CATCGCGG ACATGCCGC GTAATTCCGG CGGCCACAC CACCAATCG CGTGCACGT GCGATGTAA CGGTCGGGG ATCGGGGG AGGAAAGCGA  
2401 TTCTCCCTT CCTTCTCGC CACGTTGCC GGCTTCCCC GTCAAGCTT AAATGGGG CTCCTTGTGATT TAGTGTAA CGGCACCTCG  
AAGAAGGGAA GGAAGAGCG GTGCAAGCGG CCGAAAGGG CAGTTCGAGA TTAGCCCC GAGGAATC CCAAGGCTAA ATCAGGAAT GCGTGGAGC  
2501 ACCCCAAAAA ACTTGATTG GTGTGATGGT CACGTTGG GCCATGCC TGATGACGG TTTTCGCCC TTGACGTT GAGTCACGT TCTTAATAG  
TGGGGTTTT TGAACAAAC CCACTACCA GTGCATCACC CGTAGCGG ACTATCTGCC AAAAGGGG AACTGCAAC CTCAGGTGCA AGAAATTC  
ACCTGAGAAC AAGGTTTGAC CTGTTGIGA GTGGGGATAG AGCCGATAA GAAACTAA TATTCCCAA AACGGTAAAGA GCGGATAAC CAATTGTA  
2601 TGGACTCTG TTCCAAC TGAAACACT CAACCTATC TCGGGCTATT CTTTGATT ATAAGGGATT TTGCGATT CGCCCTATG GTTAAATAT  
ACCTGAGAAC AAGGTTTGAC CTGTTGIGA GTGGGGATAG AGCCGATAA GAAACTAA TATTCCCAA AACGGTAAAGA GCGGATAAC CAATTGTA  
2701 GAGCTGATT AACAAATT TAACGGAT TTAACAAA TATTAACGTT TACAATTAA TGTTGACTC TCACTACAT CTGCTCTGAT GCCGATAGT  
CTCGACTAA TTGTTTTAA ATTGGCTTA AAATGTTT ATAATGCAA ATGTTAAAT ACCACGTGAG AGTCATGTA GACGGACTA CGGCGTATA  
~2801 TAGGCCAATC CGCTATCGC TACGGTACTG GTCATGGCT GCGCCCGAC ACCCGCCAC ACCCGCTGAC GCGCCCTGAC GGCTGTCT GCTCCGGCA  
ATTCGGTGA GGGGATAGCG ATGCACTGAC CCAGTACCGA CGGGGGCTG TGGGGGTGTG TGGGGACTG CGGGGACTG CCCGACAGA CGAGGGCGT  
2901 TCCGCTTACA GACAAGCTGT GACCGCTCC GGGAGCTGCA TGTGTAGAG GTTTCAACCG TCATCACCGA AACGGGGAG GCAGTATCT TGAAGAGGAA  
AGGCGAATGT CTGTTGACA CTGGCAGAGG CCCTCGACGT ACACAGTCTC CAAAGTGGC AGTAGGGCT TGGGGCTC CGTCATAGA ACTCTGCTT  
3001 AGGGCCTGCT GATAGCCTA TTTTATAGG TTAATGTCAT GATAATAATG GTTCTTAGA CGTCAGGTGG CACTTTGCG GGAATGTGC GGGGAACCC  
TCCCGGAGCA CTAGGGAT AAAATATCC AATTACAGTA CTATTATAC CAAAGAATCT GCAGTCACCC GTGAAAGCC CTRTACAGG CGCCTGGGG

Figure 5C

3101 TATTGTTA TTTCTAA TACATCAA TATGTATCCG CTCATGAGAC AATAACCTG ATAATGCTT CAATAATT GAAAAGGAA GAGTATGAGT  
 ATAAACAAT AAAAGATT ATGTAAGTT ATACATAGGC GAGTACTCTG TTATGGAC TATTAAGGAA GTTATTATAA CTTTTCCCTT CTCACTACTCA  
 3201 ATCAAACATT TCCGTGCGC CCTTATTCCC TTTTGCGG CATTTGCT TCCTGTTT GCTCACCCAG AACGCTGGT GAAAGTAAA GATGCTGAAG  
 TAAGTGTAA AGGCACAGCG GGAATAAGG AAAAACGCC GTAAACGGA AGGAAAAAA CGAGTGGTC TTGCGACCA CTTTCATTT CTACGACTTC  
 3301 ATCACTGGG TGCACGACTG GGTACATCG AACTGGATCT CAACAGCGGT AAGATCCTG AGAGTTTCG CCCCAGAA CGTTTCCAA TGATGAGCAC  
 TAGTCAACCC ACGTGCTCAC CCATGTAGC TTGACTAGA GTTGTGCGCA TTCTAGGAAC TCTCAAAGC GGGCTCTT GCAAAGGT ACTACTCGTG  
 3401 TTTAAAGT CTGCTATGTG GCCGGTTT GCATCTTAG GATGGCATGA CAGTAAGA ATTATGCACT GCTGCCATAA CCATGAGTGA TAACACTGCG  
 AAATTCAA GACGATACAC CGCGCCATAA TAGGGCACTA CTGCGGCCG TTCTCGTGA GCGAGGCG TATGTATAA GAGTCTACT GAAACAACT  
 3501 TACTCACCAG TCACAGAAA GCATCTTAG GATGGCATGA CAGTAAGA ATTATGCACT GTCATTCTCT TAATACGTCA CGACGGTATT GGTACTCACT ATGTCACGC CGGTGAATG  
 ATGAGTGGTC AGTGTCTTT CGTAGAATGC CTACCGTACT GTCATTCTCT TAATACGTCA CGACGGTATT GGTACTCACT ATGTCACGC CGGTGAATG  
 3601 TTCTGACAAAC GATCGGAGGA CGGAAGGAGC TAACCGCTT TTGCAACAC ATGGGGATC ATGTAACCTG CCTTGATCGT TGGGAACCGG AGCTGAATGA  
 AAGACTGTG CTAGCCTCCT GGCTCCTCG ATGGCGAA AACGTTGCG ATTGGCGAA AACGTTGCG TACATGAGC GGAACTAGCA ACCCTGGCC TCGACTTACT  
 3701 AGCCATACCA AACGACGAGC GTGACACAC GATGCCAGCA GCAATGCAA AACGTTGCG CAAACTATA ACTGGCGAAC TACTACTCT AGCTTCCCCG  
 TCGGTATGGT TTGCTGCTCG CACTGTTGCG CTACGGTGT CGTACCGTT GTGCAACGC GTTGTATAAT TGACCGCTG ATGAATGAGA TCGAAGGGCC  
 3801 CAAACAATAA TAGACTGGAT GGAGGGGAT AAAGTGCAG GACCACTCT GGCCTCGGCC CTTCCGCTG GTTGTATAAT TGCTGATAA TCTGGAGGCC  
 GTTGTATAAT ATCTGACCTA CCTCCGCCAA TTTCACGTC CTGGGAAGA CGCGAGGCC GAAGGGCGAC CGACCAATA AGCACTATT AGACCTGGC  
 3901 GTGAGCGTGC GTCTCGCGGT ATCATTGCG CACTGGGCC AGATGCTAAG CCCCTCCGTA TCGTAGTTAT CTACACGAGC GGGAGTCAGG CAACTATGAA  
 CACTCGCACC CAGAGCGCCA TAGTAACGTC GTGACCCCCG TCTACATTC GGGAGGGCAT AGCATCAA GATGTCGTC CCCTCAGTC GTTGTATACT  
 4001 TGAACGAAAT AGACAGATCG CTGAGATGG TCCCTCACTG ATTAAGCATT GGTAACTGTC AGACCAAGT TACTCATATA TACTTAGAT TGATTAAGA  
 ATCTGCTTA TCTGCTTAGC GACTCTATCC ACGGAGTGC TAATCGTAA CCATGACAG TCTGTTCA ATGAGTATAAT ATGAATCTA ACTAAATT  
 4101 CTCATTTT AATTAAAG GATCTAGGTG AAGATCCTT TTGATAACT CATGACCAA ATCCCTAAC GTGAGTTTC GTTCCACTGA GCGTCAGAAC  
 GAGTAAGAA TAAATTTTC CTAGATCCAC TTCTAGGAA AACTATTAGA GTACTGGTT TAGGAAATG CACTCAAAG CAAAGTACT GCGAGTCTGG  
 4201 CGTAGAAA GATCAAAGA TCTCTTGAG ATCCTTTT TTGATAACT CATGACCAA ATCCCTAAC GTGAGTTTC GTTCCACTGA GCGTCAGAAC  
 GGCATCTTT CTAGTTCCCT AGAAGAACTC TAGGAAAGA AGACGCCAT TAGGAGGA ACGTTGTT TTGTTGGGC GATGGTCGCC ACCAAACAAA  
 4301 GCCGGATCAA GAGCTACCA CTCCTTTCC GAAGGTAAC GGCTTCAGCA GAGGCGAGAT ACCAAACT GTCTCTAG TCTAGCCGTA GTTAGGCCAC  
 CGGCCTAGTT CTCGATGGTT GAGAAAGG CTTCCATGA CGGAAGTCGT CTGGCGCTA TGGTTATGA CAGGAAGATC ACATGGCAT CAATCGGTG

**Figure 5D**

4401 CACTTCAAGA ACTCTGTAGC ACCGCCTACA TACCTCGCTC TGCTAATCCT GTTACCACTG GCTGCTCCA GTGGCGATA GTCGTCTT ACCGGTTGG  
 GTGAAAGTCT TGAGACATCG TGGCGGATGT ATGGAGCGAG ACGATTAGGA CAATGGTCAC CGACGACGGT CACCGTATT CAGCACAGA TGGCCCAACC  
 4501 ACTCAAGACG ATAGT'TACCG GATAAGGGCG AGCGGTCGGG CTGAACGGGG GGTTCGTCA CACAGCCAG CTGGAGCGA ACGACCTACA CGAAGCTGAG  
 TGAGTCTGCC TATCAATGCC CTATTCGGG TCGCCAGCCC GACTTGCCCC CCAAGCACGT GTGTCGGTC GAACCTCGCT TGCTGGATGT GGCTGACTC  
 4601 ATACCTACAG CGTGAGCATT GAGAAAGGCC CACGCTCCC GAAGGGAGAA AGGGGACAG GTATCCGTA AGGGCAGGG TCGAACAGG AGAGGCACG  
 TATGGATGTC GCACTCGTAA CTCTTICGCG GTGCGAAGGG TCCCGCTCT TCCGCTGTC CATAGGCCAT TCGCCGTCCC AGCCTTGTC TCTCGCGTC  
 4701 AGGGAGCTC CAGGGGAAA CGCCTGGTAT CTTATAGTC CTGTCGGTT TCGCACCTC TGACTGAGC GTCGATTGT GTGATGCTCG TCAGGGGGC  
 TCCCTCGAAG GTCCCCCTT GCGGACCAT AATATCAG GACAGCCAA AGGGGGAG ACTGAACCTCG CAGCTAAAA CACTACGAGC AGTCCCCCG  
 4801 GGGAGCTATG GAAAACGCC AGCAACCGGG CCTTTTACG GTTCTGCCC TTTGCTGCC CTTTGCTCA CATGTTCTR CCTGCGTTAT CCCCTGATC  
 CCTCGGATAC CTITTGCGG TCGTGCAGG GAAAATGC CAAGGCCAA AAACGACCG GAAACAGT GTACAAGAA GGACGCAATA GGGACTAAG  
 4901 TGTGGATAAC CGTATTACCG CTTTGAGTG AGCTGATACC GCTCGCCCA GCGAACGAC CGAGCGAGC GAGTCAGTA GCGAGGAAGC GAAAGAGCC  
 ACACCTATTG GCATAATGGC GAAACTCAC TCGACTATGG CGAGCGGT CGGCTTGCTG GCTCGCGTC CTCAGTCAC CGCTCCTCG CCTCTCGCG  
 5001 CCAATACGCA AACCGCCTC CCGGGCGT TGGCGATTC ATTAAATCAG CTGGCACAG AGGTTTCCCG ACTGGAAAGC GGGCAAGTGG CGCAACGCCA  
 GGTATGCGT TTGGGGAGA GGGGCCGCA ACCGGCTAAG TAATTAGTC GACCGTCTG TCCAAAGGGC TGACCTTCG CCCGTCATC GCGTTGCGT  
 5101 TTAATGTGAG TTACCTCACT CATTAGGCAC CCCAGGCTT ACACTTATG CTTCCGGCTC GTATGTGAG TGAAATGGT AGGGATAAC AATTACAC  
 AATTACACTC AATGGAGTGA GTAATCCGTG GGGTCCGAA TGTGAATAAC GAAGGGGAG CATAACAAC ACTTAAACAC TGGCTTATG TTAATGTGAG  
 5201 AGGAACAGC TATGACCATG ATTACGAATT AATCGAGCT CGCCCGACAT TGATTATGA CTAGTTATA ATAGTAATCA ATTACGGGT CATTAGTCA  
 TCTTTGTCG ATACTGGTAC TAATGCTAA TTAAGCTCGA GCGGGCTGTA ACTAATACT GATCAATAAT TATCATTAGT TAATGCCCA GTATCAAGT  
 from pPMLCMV beginning to HindII, enhancers and promoter

501 TAGCCCCATAT ATGGAGTCC GCGTTACATA ACTTACGGTA AATGGCCCG CTTGGTGTGAC GCCAACGAC CCCCCCCAT TGACGTCAT ATGACGGTAT  
 ATCGGGTATA TACCTCAAG CGCAATGTAT TGAATGCCAT TTACCGGGCG GACCGACTGG CGGGTTCTG GGGGGGGTA ACTGCAAGTA TTAATGCA  
 5401 GTTCCCCATAG TAACGCCAT AGGGACTTC CATTGACGTC AATGGGGTA GTATTACGG TAAACTGCC ACTTGGCAGT ACATCAAGTG TATCATATGC  
 CAAGGGTATC ATGGGGTAA TCCCTGAAAG GTAATGCGAG TTACCCACT CATAATGCC ATTGACGGG TGAACCGTCA TGTAGTTCAC ATAGTATACG  
 5501 CAAGTACGCC CCCTATGAC GTCAATGACG GTAATGGCC CCCCTGGCAT TATGCCAGT ACATGACCTT ATGGGACTT CCTACTTGGC AGTACATCTA  
 GTTCATGCGG GGGATAACTG CAGTACTGCA CATTACCG GCGGACCGTA ATACGGTCA TGTACTGGAA TACCCCTGAA GGATGAAACCG TCATGTAGAT

**Figure 5E**

5601 CGTATTAGTC ATCGCTATTA CATGGTGAT GCGGTTTGG CAGTACATCA ATGGCGTGG ATAGCGTT GACTCACGGG GATTCCAAG TCTCCACCC  
GCATAATCAG TAGCGATAAT GGTACCACTA CGCCAAACC GTCATGTAGT TACCCGACC TATGCCAAA CTGAGTGCC AGAGGTTTC AGAGGTTGGG  
5701 ATGGACGTCA ATGGGAGTT GTTGGCAC CAAATCACG GGGACTTCC AAAATGCTGT AACAACTCCCG CCCCCATGAC GCAAATGGGC GGTAGGGCGTG  
TAACGTGAGT TACCTCAA CAAACCGTG GTTTAGTG CCCTGAAGG TTTACAGCA TGTTGGGC GGGTAACTG CGTTACCCG CAACTCCGAC  
5801 TACGGTGGGA GGTCTATATA AGCAGAGCTC GTTAGTGAA CCTCAGATC GCCTGGAGAC GCCATCCACG CTGTTTGAC CTCCATAGAA GACACCGGA  
ATGCCACCCCT CCAGATATAT TCGTCTCGAG CAAATCACTT GGCACTCTAG CGGACCTCTG CGGTAGGTGC GACAACCTG GAGGTATCTT CTGTGGCCCT  
5901 CCGATCCAGC CTCCGGGGCC GGGAACGGTG CATTGAAACG CGGATCCCC GTGCCAAGAG TGACGTAGT ACCGCTATA GAGTCATAG GCCCACCCCC  
GGCTAGGTGC GAGGGCCCGG CCCTGCCAC GTAACCTTGC GCCTAAGGGG CACGGTTCTC ACTGCATCA TGGCGGATAT CTCAGATAC CGGGGGGG  
6001 TTGGCTCGTT AGAACGGGGC TACAATAAT ACATAACCT ATGTATCATA CACATACGAT TTAGGTACA CTATAGATA ACATCACTT TGCCCTTCTC  
AACCAGCAA TCTGGGCCG ATGTAATA TGTATGGAA TACATAGTAT GTGTATGCTA AATCCACTGT sp6 RNA start  
sp6 promoter  
6101 TCCACAGGTG TCCACTCCA GGTCCAATG CAGGCATGG CGGCCATCGA TT (SEQ ID NO. 25)  
AGGTCCAC AGGTGAGGGT CGAGGTGAC GTCCGGTAC GCGGGTAGCT AA cloning linker

**Figure 5F**